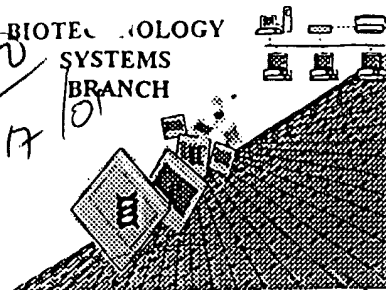


RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH

7500
10.17.01

0420/0360



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/682,769

Source: OIPE

Date Processed by STIC: 10/18/2001

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>



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CRF
OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/682,769

DATE: 10/18/2001

TIME: 06:50:00

Input Set : C:\Crf3\Datahold\EFS\09682769\PTO_MS.txt

Output Set: N:\CRF3\10182001\I682769.raw

1 <110> APPLICANT: Monsanto Co Rangwala,
W--> 2 <120> TITLE OF INVENTION: Cotton Event PV-GHGT07(1445) and Compositions and
W--> 3 <130> FILE REFERENCE: 38-21(52249)<150> 60/243190<151> 2000-10-
C--> 4 <140> CURRENT APPLICATION NUMBER: US/09/682,769
C--> 4 <141> CURRENT FILING DATE: 2001-10-17
E--> 4 <160> NUMBER OF SEQ ID: 12 <170> PatentIn version 3.0<210> 1<211> 25<212> DNA<213>

ERRORED SEQUENCES

Does Not Comply
Corrected Diskette Needed

see print out of the sequence listing
attached to this report. PatentIn failed
to save the file in the correct format.
Please contact Bob Wax at 703-308-4214
for help in getting file in correct format.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/682,769

DATE: 10/18/2001

TIME: 06:50:01

Input Set : C:\Crf3\Datahold\EFS\09682769\PTO_MS.txt

Output Set: N:\CRF3\10182001\I682769.raw

L:2 M:283 W: Missing Blank Line separator, <120> field identifier
L:3 M:283 W: Missing Blank Line separator, <130> field identifier
L:4 M:270 C: Current Application Number differs, Replaced Current Application No
L:4 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:4 M:283 W: Missing Blank Line separator, <160> field identifier
L:5 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:5 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:5 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:5 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:5 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQ ID NO
L:5 M:283 W: Missing Blank Line separator, <400> field identifier
L:24 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:24 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:24 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:31 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:31 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:31 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:38 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:38 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:38 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:55 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:55 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:55 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:78 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:78 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:78 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:85 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:85 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:85 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:92 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:92 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:92 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:99 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:99 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:99 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:4 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (12) Counted (9)